STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/757,077A_

Source: _IFW16_

Date Processed by STIC: _11/09/2006_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

RAW SEQUENCE LISTING   DATE: 11/09/2006
PATENT APPLICATION: US/10/757,077A   TIME: 13:13:43

Input Set : **N:\efs\10757077a_efs\17355CIP3SeqList.txt**
Output Set: **N:\CRF4\11092006\J757077A.raw**

```
  4 <110> APPLICANT: Steward, Lance E.
  5         Fernandez-Salas, Ester
  6         Herrington, Todd
  7         Aoki, Kei Roger
  9 <120> TITLE OF INVENTION: Clostridial Neurotoxin Compositions and
 10         Modified Clostridial Neurotoxins
 13 <130> FILE REFERENCE: 17355CIP3 (BOT)
 15 <140> CURRENT APPLICATION NUMBER: US 10/757,077A
 16 <141> CURRENT FILING DATE: 2004-01-14
 18 <150> PRIOR APPLICATION NUMBER: US 09/910,346
 19 <151> PRIOR FILING DATE: 2001-07-20
 21 <150> PRIOR APPLICATION NUMBER: US 09/620,840
 22 <151> PRIOR FILING DATE: 2000-07-21
 24 <150> PRIOR APPLICATION NUMBER: US 10/163,106
 25 <151> PRIOR FILING DATE: 2003-06-04
 27 <160> NUMBER OF SEQ ID NOS: 38
 29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
```

Does Not Comply
Corrected Diskette Needed
(pg -1)

ERRORED SEQUENCES

E--> 555 <210> SEQ ID NO: (33) 32

If Intentionally skipped sequence
Pls use this format

<210 — 000
<4007 — 000
See pg-2 for Error Explanation

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 11/09/2006
PATENT APPLICATION: US/10/757,077A   TIME: 13:13:44

Input Set : **N:\efs\10757077a_efs\17355CIP3SeqList.txt**
Output Set: **N:\CRF4\11092006\J757077A.raw**

Error Explanation

<u>Skipped Sequences(NEW RULES):</u>
Sequence(s)__missing. If intentional, please use the following format for
each skipped sequence.
<210> sequence id number
<400> sequence id number
000

Seq#:32

VERIFICATION SUMMARY   DATE: 11/09/2006
PATENT APPLICATION: US/10/757,077A   TIME: 13:13:44

Input Set : **N:\efs\10757077a_efs\17355CIP3SeqList.txt**
Output Set: **N:\CRF4\11092006\J757077A.raw**

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L:205 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:209 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:228 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:232 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:251 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:255 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:274 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:278 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:297 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:301 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:320 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:324 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:343 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:347 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:366 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:370 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:555 M:214 E: (33) Seq.# missing, SEQ ID NO:32
```